

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/960,481

DATE: 10/10/2001

TIME: 08:51:06

Input Set : D:\15367C seq listing.txt

Output Set: N:\CRF3\10102001\I960481.raw

1 <110> APPLICANT: Byrum, Joseph R.
 2 Heck, Gregory R.
 3 La Rosa, Thomas J.
 5 <120> TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
 6 Plants
 8 <130> FILE REFERENCE: 38-21(15367)C
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/960,481
 C--> 10 <141> CURRENT FILING DATE: 2001-09-24
 10 <150> PRIOR APPLICATION NUMBER: US 09/306,349
 11 <151> PRIOR FILING DATE: 1999-05-10
 13 <160> NUMBER OF SEQ ID NOS: 53893
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 300
 17 <212> TYPE: DNA
 18 <213> ORGANISM: Glycine max
 20 <223> OTHER INFORMATION: Clone ID: 701036901H1
 22 <400> SEQUENCE: 1
 24 tgtaaattctc aagatcagat cctcatctac caccattcca ggtgtagcca tcaaaacaac 60
 26 gtttggttaa caaaacccat tatcagatca gatccgatca gatcctcttc agattcggat 120
 28 caccgccgag tccatgagca cgccagcgtc ttcagggggc acgggcctct tctctggatt 180
 30 cacgcgttta tgcaagggcc ttgcggtggt gcttggttgc ggtcacattg cggttcagtt 240
 32 tatcccttca gctgttaact atcttgctct tattcccgcc aggacaatcc catttgctg 300
 35 <210> SEQ ID NO: 2
 36 <211> LENGTH: 297
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Glycine max
 40 <223> OTHER INFORMATION: Clone ID: 701036902H1
 42 <400> SEQUENCE: 2
 44 aaattacata gcctcaataa tctgagagag aaacaaaaac aatgaagggt tccattgcca 60
 46 agttccactc ccttccccta tgttttttcc ttctcttggc cttcaacaca aaacccctcc 120
 48 tagcagcaga accagagcca gtggttgata agcaagggaa ccctttggag ccaggagtag 180
 50 ggtactacgt gtggccactt tgggctgatg aaggaggcct cacactagge caaacaagga 240
 52 acaagacatg ccctctttat gttatccgtg acccttcatt cattggcacc ccatttc 297
 55 <210> SEQ ID NO: 3
 56 <211> LENGTH: 295
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Glycine max
 60 <220> FEATURE:
 61 <221> NAME/KEY: unsure
 62 <222> LOCATION: (1)..(295)
 63 <223> OTHER INFORMATION: unsure at all n locations
 64 <223> OTHER INFORMATION: Clone ID: 701036903H1
 66 <400> SEQUENCE: 3
 68 gcctagccat tgatttgga aagtccaagc ttcaaagatg agtaagcttc agagtgatgc 60
 W--> 70 tcttagagaa gctatctcan gaatcatgac tgattccaag gaaaagaaca ggaaatttgt 120
 72 ggagaccatt gaactccaaa ttgggctgaa aaactatgat ccccagaagg acaagcgttt 180
 W--> 74 tantggctct gttaagttgc ctcacattcc tcgcccccaag atgaagattt gtatgcttgg 240

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76 agatgcccaa cacgttgaag aggctgagaa gataggattg gactacatgg atgtc      295
79 <210> SEQ ID NO: 4
80 <211> LENGTH: 291
81 <212> TYPE: DNA
82 <213> ORGANISM: Glycine max
84 <223> OTHER INFORMATION: Clone ID: 701036904H1
86 <400> SEQUENCE: 4
88 tttttcttct atgaacacat ggtggctgag ttgaggggaa gtacttagaa acagaaacac    60
90 agagatgaaa ccagaaacat ctctgaactt tactttgatg ctgactttct tactttatct    120
92 gaacttggtt atgggagtgc tgagcgttga tcattatcgc agagttgact ttcctgatga    180
94 attcgttttt ggttcaggca cctctgctta tcaggtggaa ggagcttcta acgaagatgg    240
96 gagaactcct agcatctggg atacctttgc ccatgcggtg tatgagcagg a          291
99 <210> SEQ ID NO: 5
100 <211> LENGTH: 292
101 <212> TYPE: DNA
102 <213> ORGANISM: Glycine max
104 <223> OTHER INFORMATION: Clone ID: 701036905H1
106 <400> SEQUENCE: 5
108 gtttttgccc accacttttg ctttttctgc tcaaattctg tgctcatcgc cctcgtcgtg    60
110 ttcactttgt cactgctaata ggcatttttg ctggggaaac ggcgctcttc gcaaaacgaa    120
112 ttggcaaaac ggcgcagaga caaatcgcca gaaggcattg aattggagga gaatcccgag    180
114 gtgttcaaaa gttctttgct gaaggactcc aaagaacaca tcatcagtac gtactttcct    240
116 gatgtgcgca cacttttcga tgctttccaa cgcggtctca aattgccgaa ca          292
119 <210> SEQ ID NO: 6
120 <211> LENGTH: 293
121 <212> TYPE: DNA
122 <213> ORGANISM: Glycine max
124 <220> FEATURE:
125 <221> NAME/KEY: unsure
126 <222> LOCATION: (1)..(293)
127 <223> OTHER INFORMATION: unsure at all n locations
128 <223> OTHER INFORMATION: Clone ID: 701036906H1
130 <400> SEQUENCE: 6
W--> 132 caaatttttcg gatctcttct gtttcttcga atncaaaacc ctaattcccc caatttcatt    60
134 tttcaacctt gaaaaacctt ctctctctct ctgagttccc aacgggtcatc accacagctc    120
136 gatcaacagg gacatatattg ctactgaaga gatgtctcct gcttccaaat ctaagtctaa    180
138 agacaaaaat gctggcaagg aagctcaaaa gtcttctgcc aagtcttcag gatctggtaa    240
140 tgccgtggct ggtgtaccag ctagtgcata caatccttta tcgggaacat cca          293
143 <210> SEQ ID NO: 7
144 <211> LENGTH: 292
145 <212> TYPE: DNA
146 <213> ORGANISM: Glycine max
148 <223> OTHER INFORMATION: Clone ID: 701036907H1
150 <400> SEQUENCE: 7
152 ggcactcaaa cattgtttgt ttaggaaga ggaatcttga gaaatggcag gggacagcaa    60
154 gagcaagatt ctgttcacgc gagggactgg ttacatcgga aaattcatcg tggaagccag    120
156 cgccaaagct ggccacccaa ctttcctttt ggtgagggag tccactctct ccaaccccg    180
158 aaaatcatcc ctcatcgaca atttcaaagg cctcgggtgc aatcttggtt tcggagatct    240
160 atacgatcac caaagtttgg tgagtgcata aaagcaggtc gatgtggtaa tt          292

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163 <210> SEQ ID NO: 8
164 <211> LENGTH: 86
165 <212> TYPE: DNA
166 <213> ORGANISM: Glycine max
168 <220> FEATURE:
169 <221> NAME/KEY: unsure
170 <222> LOCATION: (1)..(86)
171 <223> OTHER INFORMATION: unsure at all n locations
172 <223> OTHER INFORMATION: Clone ID: 701036908H1
174 <400> SEQUENCE: 8
W--> 176 gaaaaanaaa aaaacngaac aactacngaa antcaaaacc aacttaaatt tnttatggat 60
W--> 178 ccannttnca tnggttangt tgtttt 86
181 <210> SEQ ID NO: 9
182 <211> LENGTH: 293
183 <212> TYPE: DNA
184 <213> ORGANISM: Glycine max
186 <220> FEATURE:
187 <221> NAME/KEY: unsure
188 <222> LOCATION: (1)..(293)
189 <223> OTHER INFORMATION: unsure at all n locations
190 <223> OTHER INFORMATION: Clone ID: 701036909H1
192 <400> SEQUENCE: 9
W--> 194 tcaaaccaaa aaacacaaca aaaganagan agaaaganac aaagagttcc ttgcttttgta 60
W--> 196 gtaagatata tccatcaaaa atgggttttg cttggactag aaganacgtt ttccgccgtg 120
W--> 198 cnaagaaggn gaaggaacaa agtcaactat tagaagttga ggttggtgatt cccaaccact 180
200 ttgctgtgcc ggtgagtttg gagctgatga cagatccggt gactttatca accggcatta 240
202 cctacgaccg tgtgagcatc gagaagtgga tcgagggaga aaacagaacc tgc 293
205 <210> SEQ ID NO: 10
206 <211> LENGTH: 141
207 <212> TYPE: DNA
208 <213> ORGANISM: Glycine max
210 <220> FEATURE:
211 <221> NAME/KEY: unsure
212 <222> LOCATION: (1)..(141)
213 <223> OTHER INFORMATION: unsure at all n locations
214 <223> OTHER INFORMATION: Clone ID: 701036910H1
216 <400> SEQUENCE: 10
W--> 218 tggagtttta ntttaaacta tggctatggt aatggaaaac cgnctttttt ncgtgttttc 60
W--> 220 accactttcg tttcttgatc agagtcaccc ccaaaaaaca aggcctaant cgccttcttc 120
W--> 222 ctctttccct tctttntttt t 141
225 <210> SEQ ID NO: 11
226 <211> LENGTH: 290
227 <212> TYPE: DNA
228 <213> ORGANISM: Glycine max
230 <220> FEATURE:
231 <221> NAME/KEY: unsure
232 <222> LOCATION: (1)..(290)
233 <223> OTHER INFORMATION: unsure at all n locations
234 <223> OTHER INFORMATION: Clone ID: 701036911H1

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236 <400> SEQUENCE: 11
238 ctcccgcac tctctcgtcgc tccgccacgc cttcgcgag gccgacccgt ggacggagct 60
240 catcgaccgg agctccatgt cccgaccgga caccctcgcc gaagcatact ctcggatccg 120
W--> 242 taagaacttc gcctanttcc gcgtcaatta cctaacccta atcgtgttgg cgctcgcggt 180
244 ctcgctcatc acgcatccgt tctcgctatt cgtcctcttc ggtctcctcg cgctcggtgc 240
W--> 246 gttcctttat ctctttcgcc cttcggtatca gcctgtcgtc ntcttcggac 290
249 <210> SEQ ID NO: 12
250 <211> LENGTH: 285
251 <212> TYPE: DNA
252 <213> ORGANISM: Glycine max
254 <220> FEATURE:
255 <221> NAME/KEY: unsure
256 <222> LOCATION: (1)..(285)
257 <223> OTHER INFORMATION: unsure at all n locations
258 <223> OTHER INFORMATION: Clone ID: 701036912H1
260 <400> SEQUENCE: 12
W--> 262 ggtgacgaga gccaccaccc accaccgcaa gcgcaantga aaaggttact gaagttggat 60
W--> 264 ggtacnttct tggcgaggat caacaacaga ttggccctta cgccttctct gagttgtgcc 120
266 agcatttctt gaatggatac ctctccgaaa atacgtttgt gtggtctgaa ggaagtagtg 180
268 agtggcagcc cttgtcctcg gtttctgacc tgtgggctca aattaaccgg caggacctga 240
270 ttcttcaact actgtgtctg ctctgatgt tgatgagttt gaaag 285
273 <210> SEQ ID NO: 13
274 <211> LENGTH: 289
275 <212> TYPE: DNA
276 <213> ORGANISM: Glycine max
278 <220> FEATURE:
279 <221> NAME/KEY: unsure
280 <222> LOCATION: (1)..(289)
281 <223> OTHER INFORMATION: unsure at all n locations
282 <223> OTHER INFORMATION: Clone ID: 701036913H1
284 <400> SEQUENCE: 13
286 cacttcttga accgtgcctt aactaaacta gagctagaat tagagttagc taccttgcct 60
W--> 288 aattcacaaa cgcgtccctc tacggctcta cctattagct atcttntttg tgctgtgatt 120
290 gaaattaatt tgtgatagct caccatggcc attgctactt gtgatcatga taagaacgca 180
W--> 292 aaggcctttgc agttcattga ggatattgacc caaacactg aaagtgtcca aganagggtc 240
W--> 294 cttgctgaga ttctcagcca aaactcccag accgaatacc tganacggt 289
297 <210> SEQ ID NO: 14
298 <211> LENGTH: 282
299 <212> TYPE: DNA
300 <213> ORGANISM: Glycine max
302 <220> FEATURE:
303 <221> NAME/KEY: unsure
304 <222> LOCATION: (1)..(282)
305 <223> OTHER INFORMATION: unsure at all n locations
306 <223> OTHER INFORMATION: Clone ID: 701036914H1
308 <400> SEQUENCE: 14
W--> 310 actgatcata aatcacatag ctagtgtttt aaagtgagag anaaagaaag aagaanatgg 60
312 caaactttcc agttgttgac atggggaaac ttaacaccga agagagagca gcagccatgg 120
W--> 314 agattataaa anatgcctgt gagaactggg gtttctttga gttggtgaac catgggatat 180

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316 ccattgagtt gatggacact gtggagagggc tcacaaaaga gcactacaag aagactatgg 240
318 agcaaagggtt caaagaaatg gtggccagca aagggttga gt 282
321 <210> SEQ ID NO: 15
322 <211> LENGTH: 290
323 <212> TYPE: DNA
324 <213> ORGANISM: Glycine max
326 <220> FEATURE:
327 <221> NAME/KEY: unsure
328 <222> LOCATION: (1)..(290)
329 <223> OTHER INFORMATION: unsure at all n locations
330 <223> OTHER INFORMATION: Clone ID: 701036915H1
332 <400> SEQUENCE: 15
334 ctaaagcaaa cccagttgtc aagggtgaa aagcttgata agatctaaga tttgaagttg 60
336 aagttccatg ccctgagatg aagttatgca caatttcac atgaaatgat ccgagtttcc 120
W--> 338 ttcacatttt attggtgttc ttattatatt gttatatgtg gtgggtgact tgagtnattg 180
340 gacatctctg caatttgaag gactctttca tgtgattggt gaaaatatat atgttggcca 240
342 ctgttggttt taagtgcggt ttggtgaaag ttggttggtg ttgttagctt 290
345 <210> SEQ ID NO: 16
346 <211> LENGTH: 170
347 <212> TYPE: DNA
348 <213> ORGANISM: Glycine max
350 <223> OTHER INFORMATION: Clone ID: 701036916H1
352 <400> SEQUENCE: 16
354 gtttctctct acaagtccat atgtaaaatg gataacagaa agaaaaagaa aaatcacttt 60
356 ggtttgagaa ctttttaaag tttatatata ctgtgttaag gttcataaaa ctgtagactg 120
358 atttgtgtga catgctccac agaaccttaa attttcctct attttgtcct 170
361 <210> SEQ ID NO: 17
362 <211> LENGTH: 302
363 <212> TYPE: DNA
364 <213> ORGANISM: Glycine max
366 <220> FEATURE:
367 <221> NAME/KEY: unsure
368 <222> LOCATION: (1)..(302)
369 <223> OTHER INFORMATION: unsure at all n locations
370 <223> OTHER INFORMATION: Clone ID: 701036917H1
372 <400> SEQUENCE: 17
W--> 374 ggccggtttcg aaagcnatga cgctatccac cgcgccctca ccgccaccgg cattagaaga 60
376 ggagccggtta cctcacgacg gcgacgatta cgacggcatc gtttcactgg acgacgaggg 120
W--> 378 tttcccgtn cgaattcccc gaatcaggaa cggcgctcga aanagcagt ttcgccgagg 180
380 attcactatc gcagacggaa ccgattgggt cttcgctggt gttgtcgctg gaagatagta 240
382 aacagagcga tttagcgcaa gattcgggga agaagatgaa gcaggccaat ctgttccaga 300
384 tt 302
387 <210> SEQ ID NO: 18
388 <211> LENGTH: 294
389 <212> TYPE: DNA
390 <213> ORGANISM: Glycine max
392 <223> OTHER INFORMATION: Clone ID: 701036918H1
394 <400> SEQUENCE: 18
396 gtttccgata tccaagggga aatggcattg gtgttcaact accaactgga aataaaggaa 60

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→ Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/960,481

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Input Set : D:\15367C seq listing.txt

Output Set: N:\CRF3\10102001\I960481.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:220 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:288 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:702 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:746 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33

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L:772 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:820 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36